Stat Comp HW 6

Jason Rights

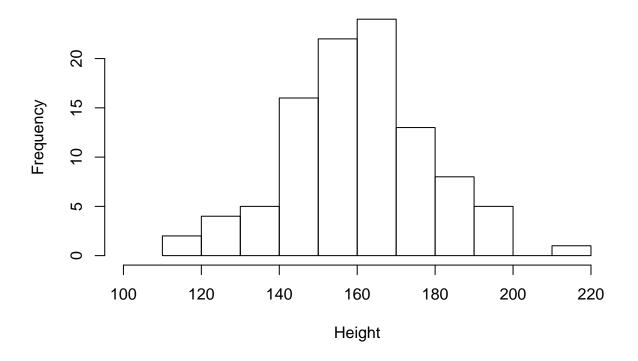
November 23, 2015

Problem 1:

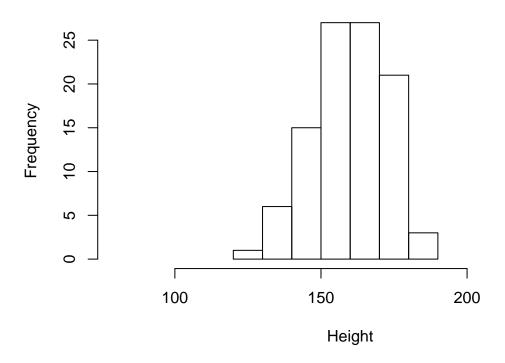
```
pop <- data.frame(m = rnorm(100,160,20), f = rnorm(100, 160, 20))

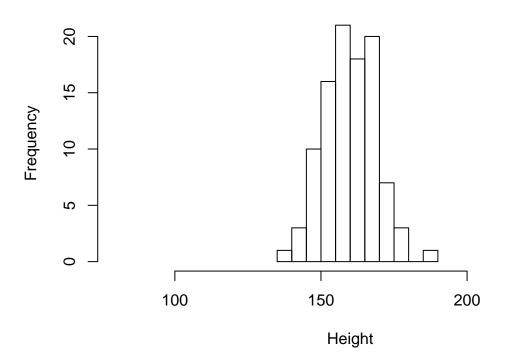
next_gen <- function(pop) {
    pop$m <- sample(pop$m)
    pop$m <- rowMeans(pop)
    pop$f <- pop$m
    pop
}

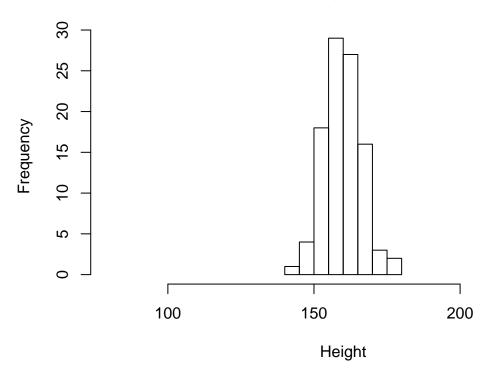
hist(pop$m,main='Male Heights Gen 1',xlab='Height',xlim=c(100,220))</pre>
```

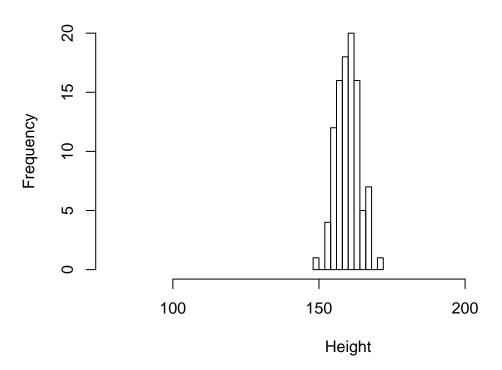


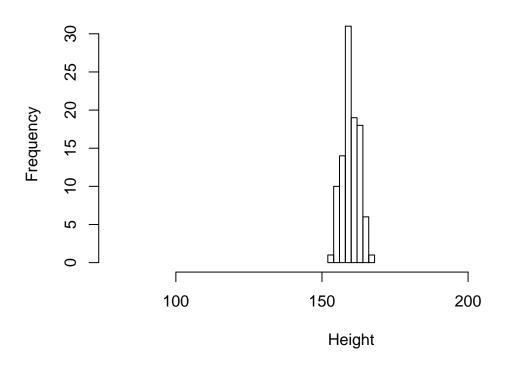
```
next.pop <- as.data.frame(matrix(NA,100,16))
colnames(next.pop) <- rep(c("m","f"),8)
for(i in c(0:7)){
  if (i==0) next.pop[,c(2*i+1):c(2*i+2)] <- next_gen(pop)
  if (i>0) next.pop[,c(2*i+1):c(2*i+2)] <- next_gen(next.pop[,c(2*i-1):c(2*i)])</pre>
```

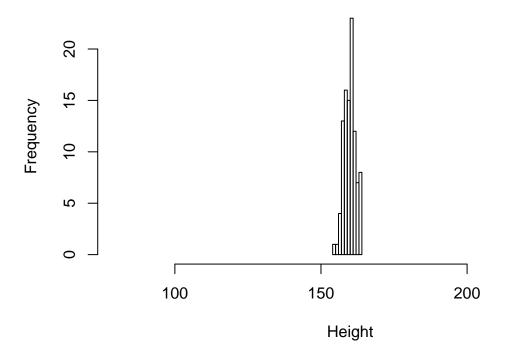


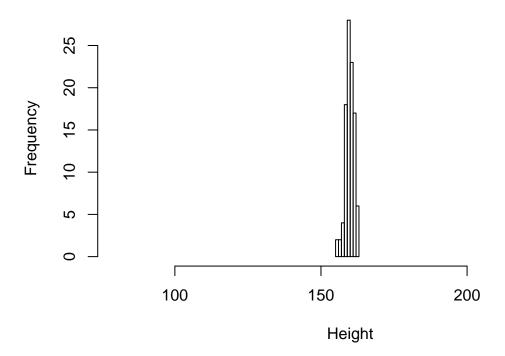


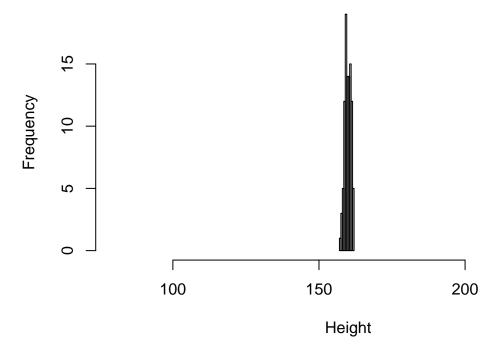












I kept the axes the same across all histograms to highlight the effect.

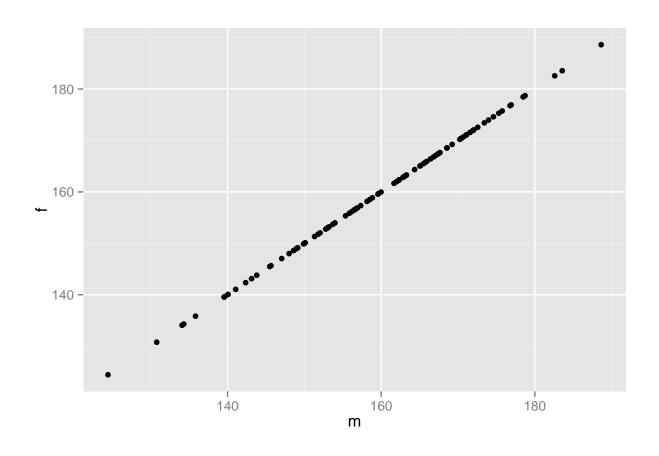
Problem 2:

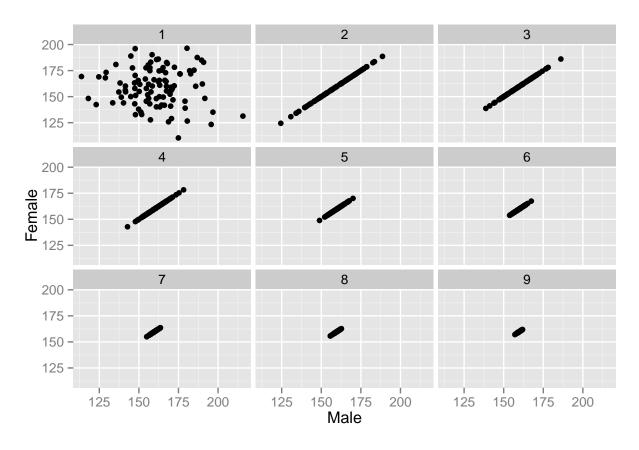
```
library(ggplot2)
total.pop <- cbind(pop,next.pop)
head(total.pop)</pre>
```

```
##
                     f
                                       f
                                                          f
                              m
                                                                   m
## 1 190.0696 162.1642 156.1841 156.1841 159.5778 159.5778 154.7926 154.7926
## 2 172.3212 160.3895 159.9736 159.9736 154.9762 154.9762 160.9827 160.9827
## 3 154.4853 177.6219 167.6430 167.6430 158.8772 158.8772 157.3395 157.3395
## 4 162.2287 164.5289 153.0110 153.0110 138.7250 138.7250 153.0829 153.0829
## 5 155.5494 151.7731 151.7752 151.7752 143.8044 143.8044 155.4974 155.4974
##
  6 180.3850 196.4489 182.5793 182.5793 177.0693 177.0693 160.7806 160.7806
##
            m
                     f
                              m
                                       f
                                                          f
## 1 157.5076 157.5076 160.9150 160.9150 160.3172 160.3172 158.9843 158.9843
## 2 161.1271 161.1271 159.9338 159.9338 160.4464 160.4464 161.9840 161.9840
## 3 159.1843 159.1843 161.9538 161.9538 162.3386 162.3386 161.7439 161.7439
## 4 154.2901 154.2901 159.5319 159.5319 160.0191 160.0191 159.6030 159.6030
## 5 157.9999 157.9999 159.9907 159.9907 158.7918 158.7918 159.4054 159.4054
## 6 164.5259 164.5259 163.4980 163.4980 161.7284 161.7284 160.0930 160.0930
##
                     f
            m
## 1 159.5691 159.5691
## 2 160.7741 160.7741
## 3 160.2483 160.2483
```

```
## 4 160.6833 160.6833
## 5 160.7252 160.7252
## 6 159.8231 159.8231
dim(total.pop)
## [1] 100 18
total.pop.male <- data.frame(stack(total.pop[,c(1,3,5,7,9,11,13,15,17)]))
total.pop.female <- data.frame(stack(total.pop[,c(2,4,6,8,10,12,14,16,18)]))
total.pop.long <- cbind(total.pop.male[,1],total.pop.female[,1])</pre>
total.pop.long <- cbind(total.pop.long,c(rep(1,100),rep(2,100),rep(3,100),rep(4,100),
colnames(total.pop.long) <- c("Male", "Female", "Gen")</pre>
head(total.pop.long)
                  Female Gen
##
            Male
## [1,] 190.0696 162.1642
## [2,] 172.3212 160.3895
## [3,] 154.4853 177.6219
## [4,] 162.2287 164.5289
## [5,] 155.5494 151.7731
## [6,] 180.3850 196.4489
```

qplot(m,f,data=next.pop)





Problem 3:

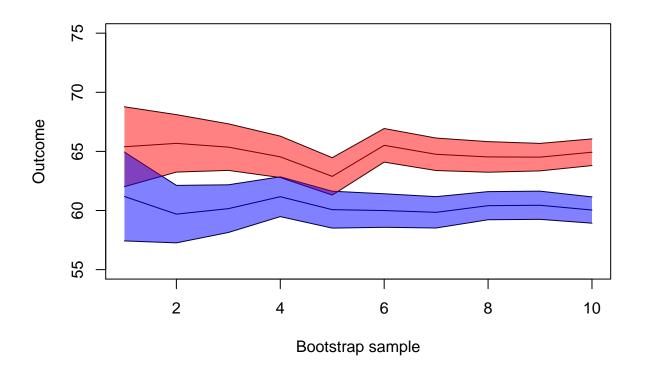
```
library(boot)
boot.ci.con <- matrix(NA,10,2)</pre>
boot.ci.treat <- matrix(NA,10,2)</pre>
for(i in seq(10)){
  N <- 250*i
  treatment <- rbinom(N,size=1,prob=.5)</pre>
  outcome <- rnorm(N,60,20)
  dat <- as.data.frame(cbind(treatment,outcome))</pre>
  for(j in seq(N)){
     dat[j,2] \leftarrow ifelse(dat[j,1] == 1, dat[j,2] + 5, dat[j,2])
  dat.con <- subset(dat, treatment==0)</pre>
dat.treat <- subset(dat, treatment==1)</pre>
  outcome.con_boot <- boot(dat.con$outcome, function(x,i) mean(x[i]), R=2000)</pre>
  outcome.treat_boot <- boot(dat.treat$outcome, function(x,i) mean(x[i]), R=2000)</pre>
  boot.ci.con[i,] <- boot.ci(outcome.con_boot)$normal[2:3]</pre>
  boot.ci.treat[i,] <- boot.ci(outcome.treat_boot)$normal[2:3]</pre>
```

```
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
```

Warning in boot.ci(outcome.con_boot): bootstrap variances needed for

studentized intervals

```
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.con_boot): bootstrap variances needed for
## studentized intervals
## Warning in boot.ci(outcome.treat_boot): bootstrap variances needed for
## studentized intervals
boot.ci.con <- cbind(boot.ci.con,NA)</pre>
boot.ci.treat <- cbind(boot.ci.treat,NA)</pre>
for(i in seq(10)){
  boot.ci.con[i,3] <- mean(c(boot.ci.con[i,1],boot.ci.con[i,2]))</pre>
  boot.ci.treat[i,3] <- mean(c(boot.ci.treat[i,1],boot.ci.treat[i,2]))</pre>
}
plot(boot.ci.con[,1],ylim=c(55,75),col="white",ylab="Outcome",xlab="Bootstrap sample")
lines(boot.ci.con[,1],ylim=c(55,80))
lines(boot.ci.con[,2])
lines(boot.ci.con[,3])
lines(boot.ci.treat[,1],ylim=c(55,80))
lines(boot.ci.treat[,2])
lines(boot.ci.treat[,3])
\#legend(100,50,legend=line,lty=c(1,1))
makeTransparent = function(..., alpha=0.5) {
  if(alpha<0 | alpha>1) stop("alpha must be between 0 and 1")
  alpha = floor(255*alpha)
  newColor = col2rgb(col=unlist(list(...)), alpha=FALSE)
  .makeTransparent = function(col, alpha) {
    rgb(red=col[1], green=col[2], blue=col[3], alpha=alpha, maxColorValue=255)
  newColor = apply(newColor, 2, .makeTransparent, alpha=alpha)
  return(newColor)
}
polygon(c(1:10,10:1),
        c(boot.ci.treat[1:10,2],boot.ci.treat[10:1,1]), col=makeTransparent('red',alpha=0.5), border = 1
polygon(c(1:10,10:1),
        c(boot.ci.con[1:10,2],boot.ci.con[10:1,1]), col=makeTransparent('blue',alpha=0.5), border = NA)
```



 $\#legend("topright", \ legend = c("Treatment", "Control"), \ col=c("red", "blue"), \ Olty=c(1:1), lwd=c(3,3), seg.$

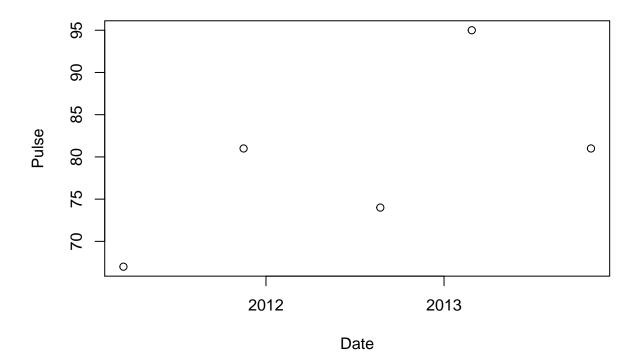
Problem 4:

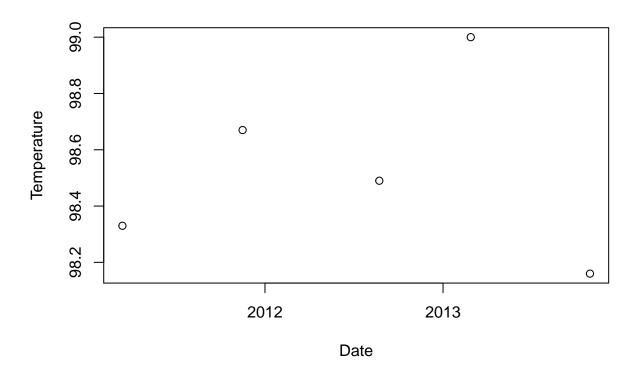
```
makePatient <- function() {</pre>
  vowel <- grep("[aeiou]", letters)</pre>
  cons <- grep("[^aeiou]", letters)</pre>
  name <- paste(sample(LETTERS[cons], 1), sample(letters[vowel], 1), sample(letters[cons], 1), sep='')</pre>
  gender <- factor(sample(0:1, 1), levels=0:1, labels=c('female', 'male'))</pre>
  dob <- as.Date(sample(7500, 1), origin="1970-01-01")</pre>
  n <- sample(6, 1)</pre>
  doa <- as.Date(sample(1500, n), origin="2010-01-01")</pre>
  pulse <- round(rnorm(n, 80, 10))</pre>
  temp <- round(rnorm(n, 98.4, 0.3), 2)
  fluid <- round(runif(n), 2)
  x <- list(name, gender, dob, doa, pulse, temp, fluid)
  names(x) <- list("name", "gender", "date_of_birth", "date_of_admission", "pulse", "temperature", "flu</pre>
  class(x) <- "medicalRecord"</pre>
  х
}
set.seed(8)
med.record <- makePatient()</pre>
med.record
```

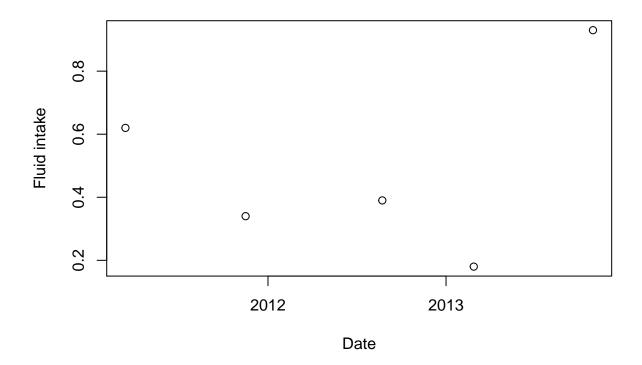
\$name

```
## [1] "Mev"
##
## $gender
## [1] male
## Levels: female male
## $date_of_birth
## [1] "1976-08-09"
## $date_of_admission
## [1] "2011-03-14" "2013-10-30" "2013-02-27" "2012-08-23" "2011-11-16"
## $pulse
## [1] 67 81 95 74 81
## $temperature
## [1] 98.33 98.16 99.00 98.49 98.67
## $fluid_intake
## [1] 0.62 0.93 0.18 0.39 0.34
## attr(,"class")
## [1] "medicalRecord"
class(med.record)
## [1] "medicalRecord"
mean.medicalRecord <- function(med){</pre>
  x <- list(mean(med$pulse),mean(med$temperature),mean(med$fluid_intake))
  names(x) <- c("mean_pulse", "mean_temperature", "mean_fluid_intake")</pre>
  return(x)
mean(med.record)
## $mean_pulse
## [1] 79.6
##
## $mean_temperature
## [1] 98.53
## $mean_fluid_intake
## [1] 0.492
print.medicalRecord <- function(med){</pre>
  x <- data.frame(med$date_of_admission,med$pulse,med$temperature,med$fluid_intake)
  x <- x[order(med$date of admission),]
  colnames(x) <- c("Date of admission", "Pulse", "Temperature", "Fluid intake")</pre>
  rownames(x) <- NULL
  plot(x[,1],x[,2],xlab="Date",ylab="Pulse")
  plot(x[,1],x[,3],xlab="Date",ylab="Temperature")
```

```
plot(x[,1],x[,4],xlab="Date",ylab="Fluid intake")
  return(x)
}
print(med.record)
```







```
##
     Date of admission Pulse Temperature Fluid intake
## 1
            2011-03-14
                            67
                                     98.33
## 2
             2011-11-16
                           81
                                     98.67
                                                    0.34
## 3
             2012-08-23
                                     98.49
                                                    0.39
                           74
             2013-02-27
## 4
                           95
                                     99.00
                                                    0.18
## 5
             2013-10-30
                                     98.16
                            81
                                                    0.93
```

```
##part 3

makePatientCohort <- function(n.patients) {
  cohort <- makePatient()
  for(i in c(2:n.patients)){
    cohort <- c(cohort,makePatient())
  }
  class(cohort) <- c("cohort")
  return(cohort)
}

x <-makePatientCohort(10)
class(x)</pre>
```

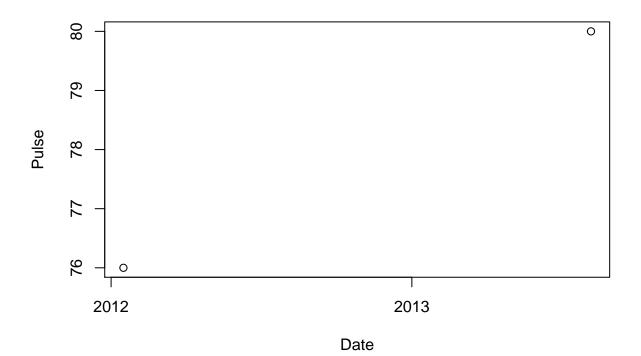
```
## [1] "cohort"
```

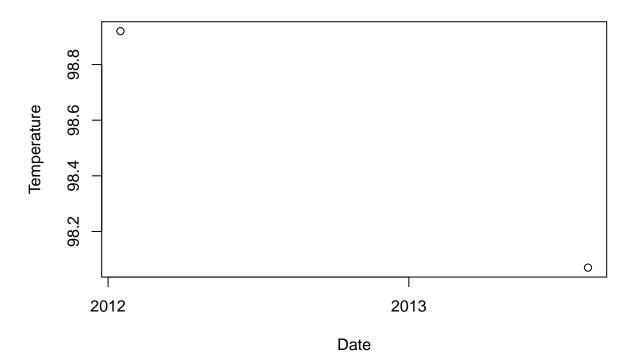
```
mean.cohort <- function(cohort){</pre>
     for(i in seq(length(cohort)/7)){
     x \leftarrow list(mean(cohort[c(7*(i-1)+1):c(7*i)] pulse), mean(cohort[c(7*(i-1)+1):c(7*i)] temperature), mean(cohort[c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1):c(7*(i-1)+1
     names(x) <- c(paste("mean_pulse_patient_",i),paste("mean_temperature_patient_",i),paste("mean_fluid_i</pre>
  print(x)
        }
}
print.cohort <- function(cohort){</pre>
     for(i in seq(length(cohort)/7)){
     x \leftarrow data.frame(cohort[c(7*(i-1)+1):c(7*i)] $\date_of_admission,cohort[c(7*(i-1)+1):c(7*i)]$ pulse,cohort[c(7*(i-1)+1):c(7*i)]$
     x \leftarrow x[order(cohort[c(7*(i-1)+1):c(7*i)]$date_of_admission),]
     colnames(x) <- c(paste("Date of admission, patient",i),"Pulse","Temperature","Fluid intake")</pre>
     rownames(x) <- NULL
     plot(x[,1],x[,2],xlab="Date",ylab="Pulse",main=paste("Patient",i))
     plot(x[,1],x[,3],xlab="Date",ylab="Temperature",main=paste("Patient",i))
     plot(x[,1],x[,4],xlab="Date",ylab="Fluid intake",main=paste("Patient",i))
     print(x)
}
mean(x)
## $`mean_pulse_patient_ 1`
## [1] 78
##
## $`mean_temperature_patient_ 1`
## [1] 98.495
##
## $`mean_fluid_intake_patient_ 1`
## [1] 0.245
## $`mean_pulse_patient_ 2`
## [1] 81.5
##
## $`mean_temperature_patient_ 2`
## [1] 98.44
##
## $`mean_fluid_intake_patient_ 2`
## [1] 0.4033333
##
## $`mean_pulse_patient_ 3`
## [1] 78
## $`mean_temperature_patient_ 3`
## [1] 98.6
## $`mean_fluid_intake_patient_ 3`
## [1] 0.65
##
## $`mean_pulse_patient_ 4`
```

```
## [1] 88.33333
##
## $`mean_temperature_patient_ 4`
## [1] 98.05
## $`mean_fluid_intake_patient_ 4`
## [1] 0.5866667
## $`mean_pulse_patient_ 5`
## [1] 83.5
## $`mean_temperature_patient_ 5`
## [1] 98.45
##
## $`mean_fluid_intake_patient_ 5`
## [1] 0.4525
##
## $`mean_pulse_patient_ 6`
## [1] 83
##
## $`mean_temperature_patient_ 6`
## [1] 98.01
##
## $`mean fluid intake patient 6`
## [1] 0.97
## $`mean_pulse_patient_ 7`
## [1] 77.5
##
## $`mean_temperature_patient_ 7`
## [1] 98.14833
##
## $`mean_fluid_intake_patient_ 7`
## [1] 0.3366667
## $`mean_pulse_patient_ 8`
## [1] 77
##
## $`mean_temperature_patient_ 8`
## [1] 98.83
## $`mean_fluid_intake_patient_ 8`
## [1] 0.445
##
## $`mean_pulse_patient_ 9`
## [1] 79.33333
## $`mean_temperature_patient_ 9`
## [1] 98.3
## $`mean_fluid_intake_patient_ 9`
## [1] 0.6583333
##
## $`mean_pulse_patient_ 10`
```

```
## [1] 81
##
## $`mean_temperature_patient_ 10`
## [1] 98.21
##
## $`mean_fluid_intake_patient_ 10`
## [1] 0.64
```

print(x)



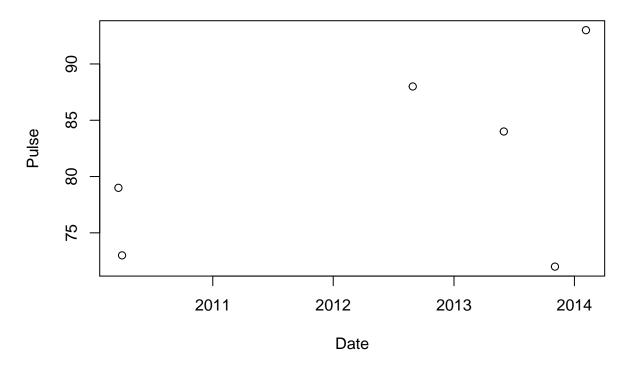




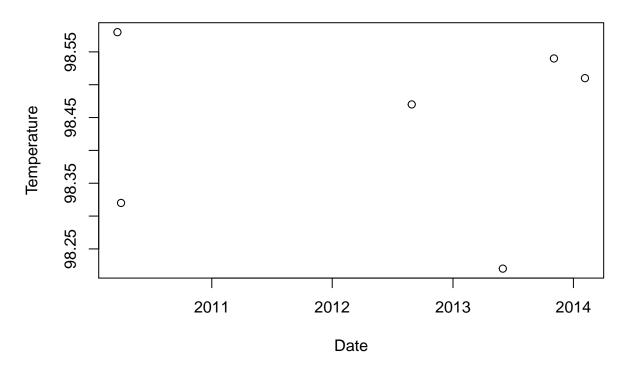


##		Date	of	admission,	patient	1	Pulse	Temperature	Fluid	intake
##	1			2	2012-01-1	6	76	98.92		0.14
##	2			2	2013-08-0	7	80	98.07		0.35

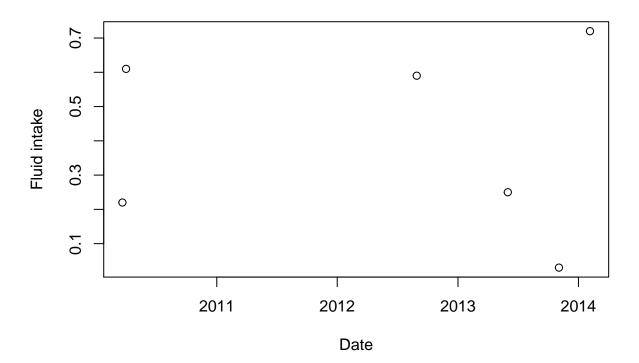
Patient 2



Patient 2

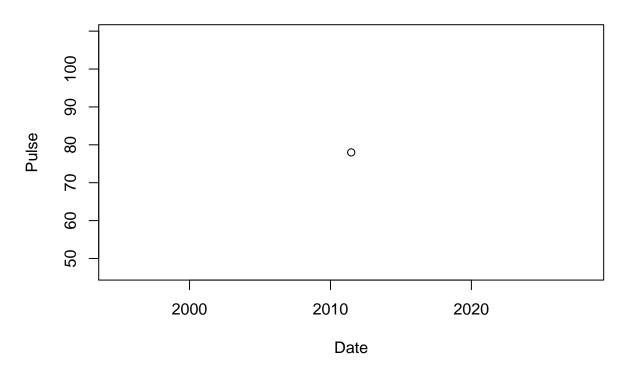


Patient 2

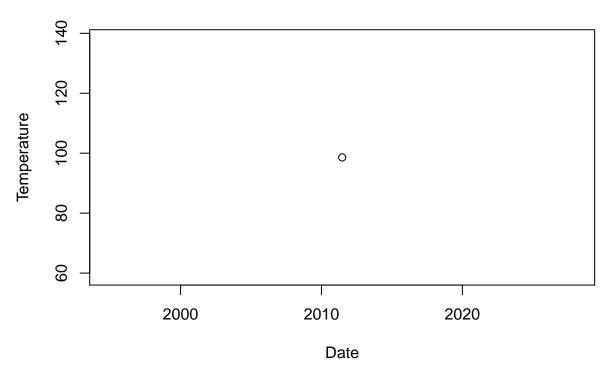


##	I	Date	of	admission,	patient	2	Pulse	Temperature	Fluid	intake
##	1				2010-03-2	21	79	98.58		0.22
##	2				2010-04-0	01	73	98.32		0.61
##	3				2012-08-2	29	88	98.47		0.59
##	4				2013-06-0	01	84	98.22		0.25
##	5				2013-11-0)3	72	98.54		0.03
##	6				2014-02-0)5	93	98.51		0.72

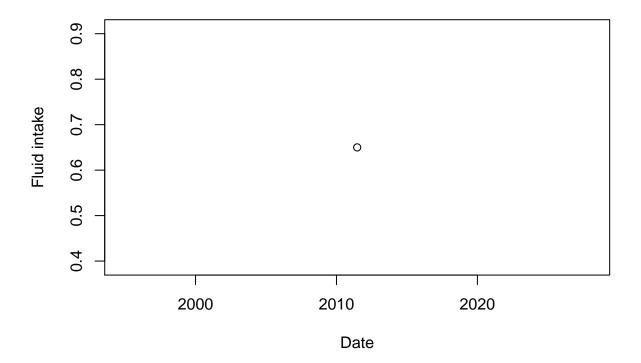
Patient 3





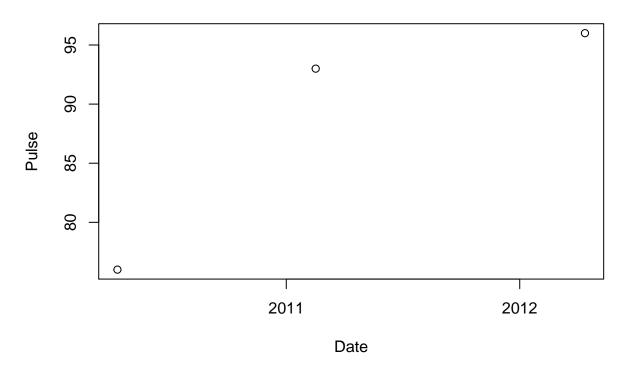


Patient 3

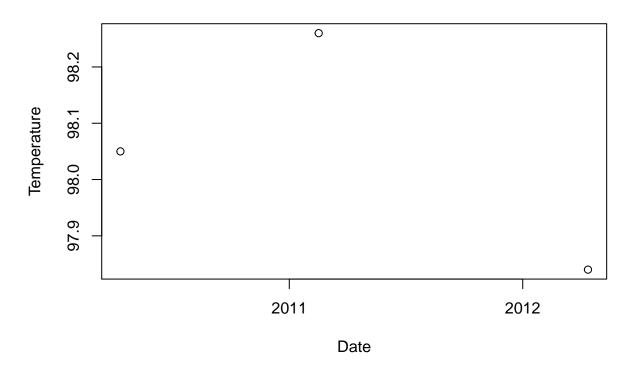


Date of admission, patient 3 Pulse Temperature Fluid intake ## 1 2011-06-22 78 98.6 0.65

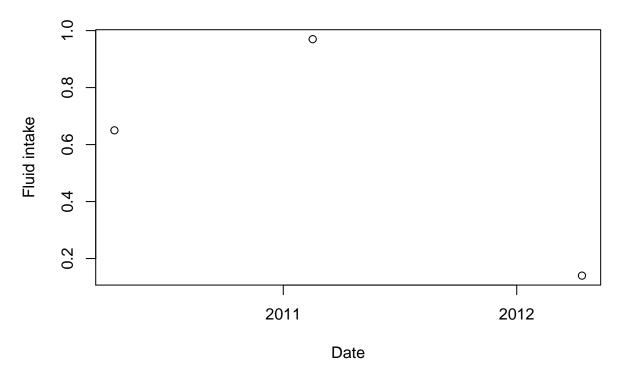




Patient 4

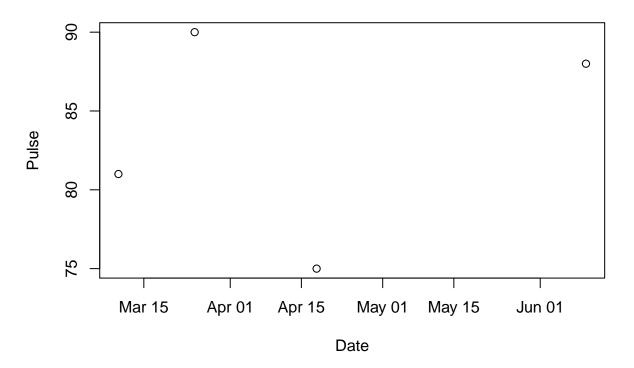


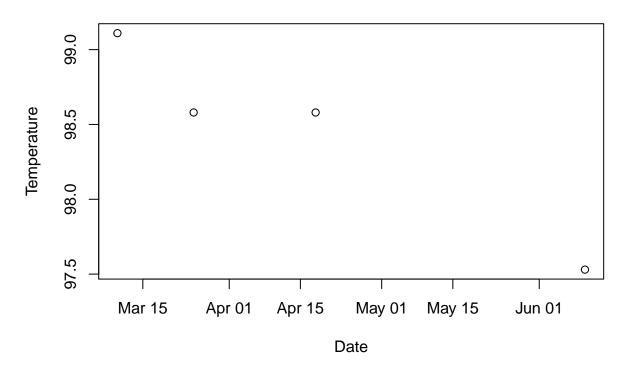




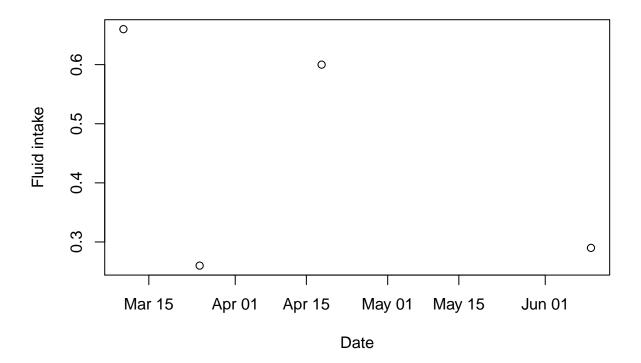
##		Date	of	admission,	patient 4	Pulse	Temperature	Fluid	intake
##	1			•	2010-04-12	2 76	98.05		0.65
##	2			•	2011-02-16	93	98.26		0.97
##	3			:	2012-04-12	96	97.84		0.14

Patient 5

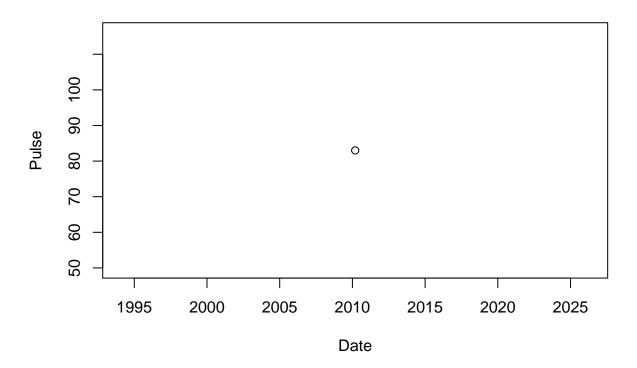




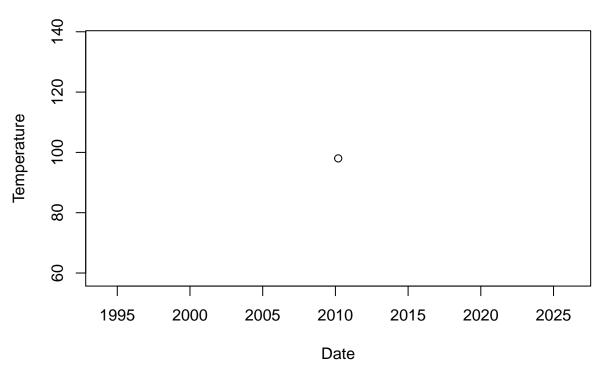
Patient 5



##		Date	of	admission, patient 5	Pulse	Temperature	Fluid intake
##	1			2010-03-10	81	99.11	0.66
##	2			2010-03-25	90	98.58	0.26
##	3			2010-04-18	75	98.58	0.60
##	4			2010-06-10	88	97.53	0.29





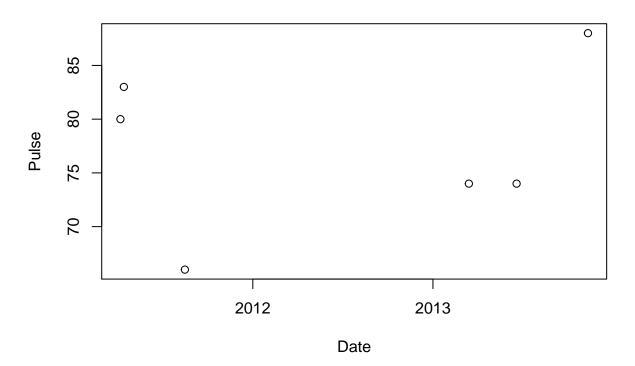




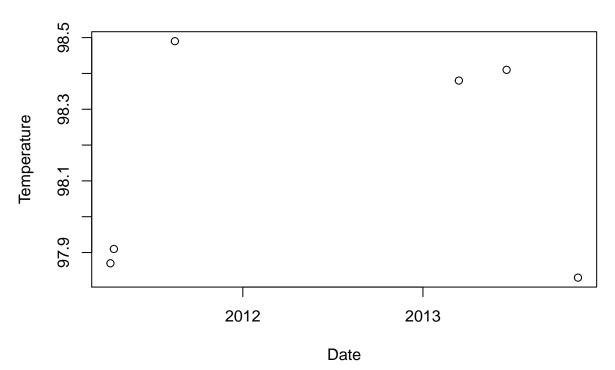


Date of admission, patient 6 Pulse Temperature Fluid intake ## 1 2010-03-12 83 98.01 0.97

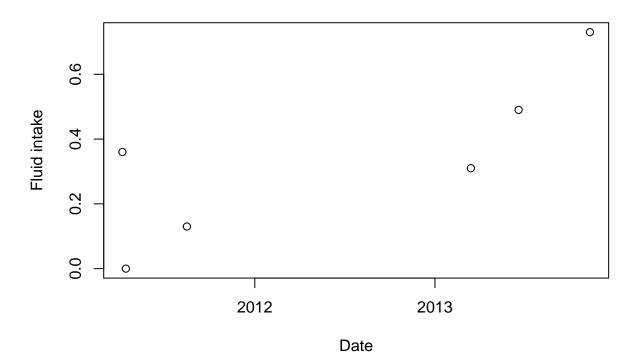
Patient 7





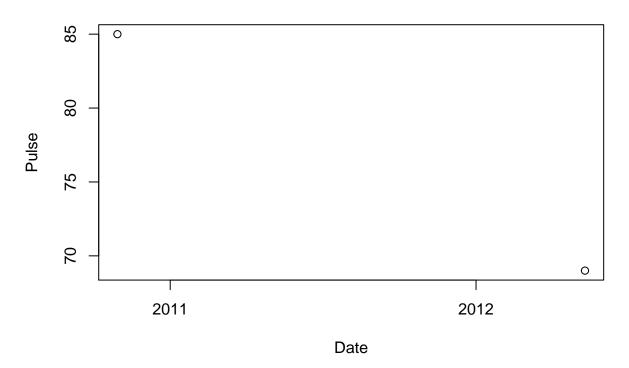


Patient 7

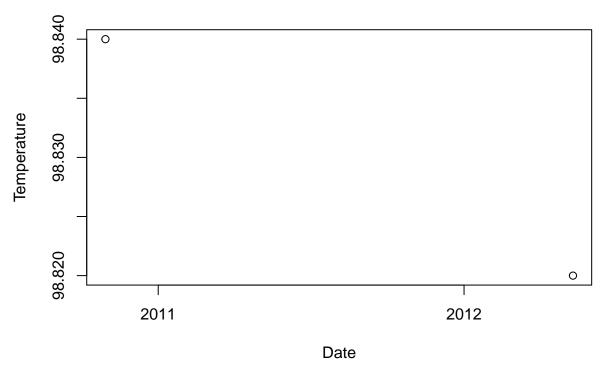


##		Date	of	admission, patient 7	Pulse	Temperature	Fluid intake
##	1			2011-04-07	80	97.87	0.36
##	2			2011-04-14	83	97.91	0.00
##	3			2011-08-16	66	98.49	0.13
##	4			2013-03-15	74	98.38	0.31
##	5			2013-06-20	74	98.41	0.49
##	6			2013-11-12	88	97.83	0.73

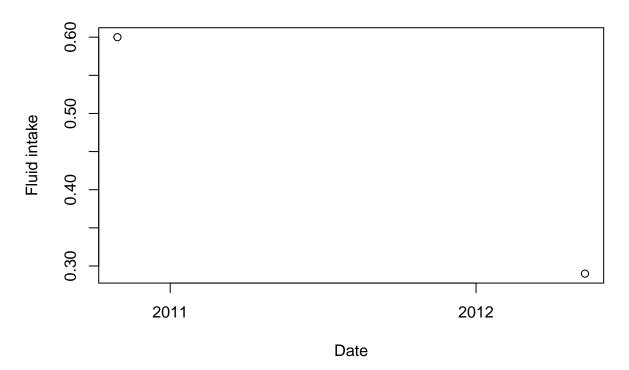
Patient 8





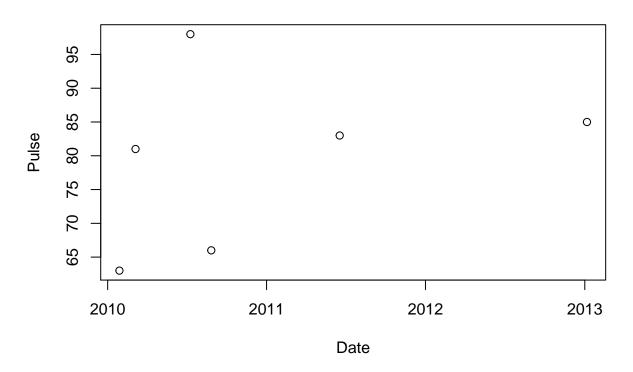


Patient 8

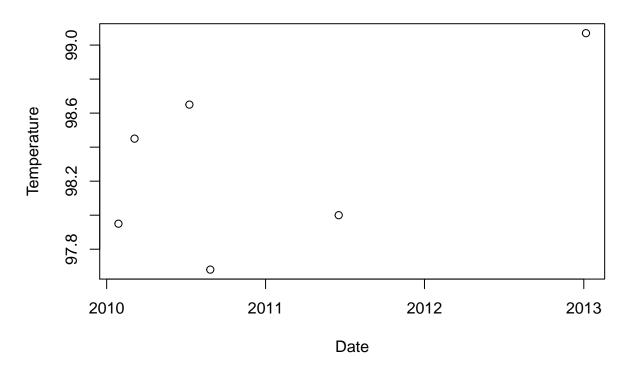


Date of admission, patient 8 Pulse Temperature Fluid intake ## 1 2010-10-30 85 98.84 0.60 ## 2 2012-05-10 69 98.82 0.29

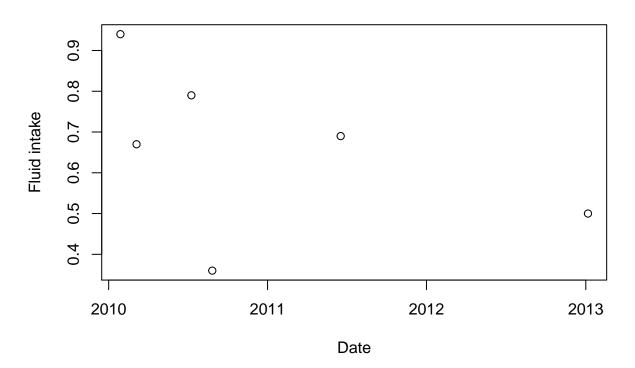
Patient 9





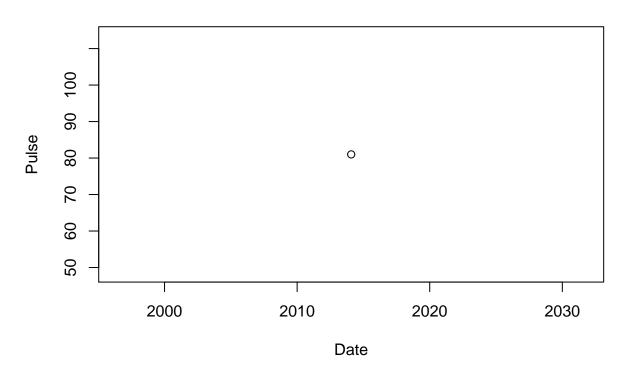


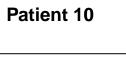
Patient 9

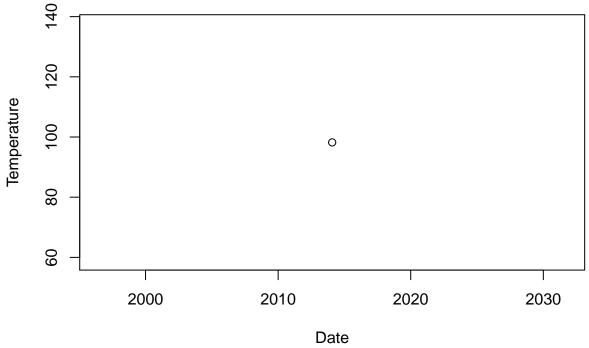


##		Date	of	admission,	patient	9	Pulse	Temperature	Fluid	intake
##	1				2010-01-2	28	63	97.95		0.94
##	2				2010-03-0)6	81	98.45		0.67
##	3				2010-07-1	0	98	98.65		0.79
##	4				2010-08-2	27	66	97.68		0.36
##	5				2011-06-1	8	83	98.00		0.69
##	6				2013-01-0)6	85	99.07		0.50

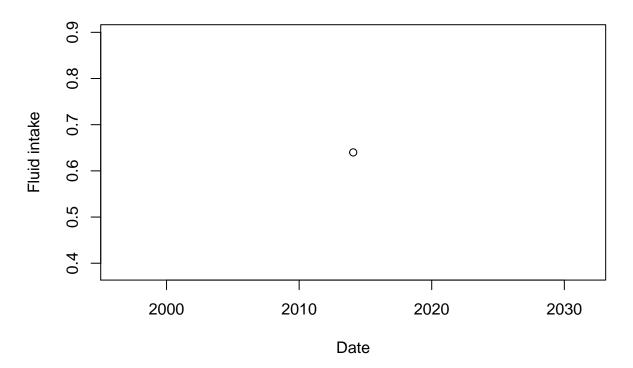
Patient 10











Date of admission, patient 10 Pulse Temperature Fluid intake ## 1 2014-01-29 81 98.21 0.64